

SEQUENCE LISTING

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<120> DATABASES OF REGULATORY SEQUENCES; METHODS OF MAKING AND USING SAME
<130> 8325-0015
<140> 09/844,501
<141> 2001-04-27
<150> 60/200,590
<151> 2000-04-28
<150> 60/214,674
<151> 2000-06-27
<150> 60/228,556
<151> 2000-08-28
<160> 24
<170> PatentIn Ver. 2.0
<210> 1
<211> 6
<212> DNA
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<223> Description of Artificial Sequence: Kpn 1 target
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ggtacc
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      oligonucleotide
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                                                                    11
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<400> 4
gcccatcact gagaaatccc ttcc
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<223> Description of Artificial Sequence: adapter
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oligonucleotide

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<211> 66
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aaaaaaaatc ttccgtgtca gctcctgaat aggatcggag acttatgaaa gttgttcaat 60
gtggga
                                                              66
<210> 9
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
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aggcacagtc gaggacttat ccta
                                                              24
<210> 10
<211> 122
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: insert
     sequence
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ceggeetegg tgttttegge ttttteetgg ecceeggeee geeaggeegg geeetetget 60
tc
                                                              122
<210> 11
<211> 249
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<400> 11
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tccgggctgg ggctgaccgg ctctgtgacc ttgggcaggt cactgcatct ctccaagcct 180
cagtttgcac gtctgtcaaa tagaggggca ttctctcact ttgcagggtc cctggaaata 240
agtgagatc
<210> 12
<211> 1042
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: accessible
     region sequence
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aaaaqqaqtt cqaqaccaqc ccggccaact ggtgaaaccc tgtctctact aaaaaaatac 120
aaaaattaqc tqqqtqtqqt qqtqcacqcc tqtcatccca qctacttqqq aqqctqaqat 180
aggaattage toggetget getgeacgee teteatecea getaettege aggetgagat 240
aggagaatcg cttgaaccca ggaggggagg cagaggttgc agtgagccga gatggcgcca 300
ctqtqaatcq cttgaaccca ggaggggagg cagaggttgc agtgagccga gatggcgcca 360
ctqtactccq qcctqqqcaa gagcaagact ccaaccaaaa aaaaaaaaaa aaagaactag 420
cagtgcccag ggctgtacac caggtgccag tactggcagc aattcttcca gttattgtga 540
taqaqccaq qqctqtacac caqqtqccaq tactqqcaqc aattcttcca qttattqtqa 600
tagattetea tgacqetaaa atacceaett tgttatttaa eeettgetaa teeacaatga 660
gttgttctca tgacgctaaa atacccactt tgttatttaa cccttgctaa tccacaatga 720
attgggcatc actttgtttt aataattctt gtatgagaag agcactcttt tccttctgat 900
agcaggcatc actttgtttt aataattctt gtatgagaag agcactcttt tccttctgat 960
agcaatgtgg ctccaactac tggctgatgt gagacggtac cggatgtggc tccaactact 1020
ggctgatgtg agacggtacc gg
                                                           1042
<210> 13
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<220>
<223> Description of Artificial Sequence: adapter
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     end
<400> 13
gatcgaattc ag
                                                           12
<210> 14
<211> 8
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<213> Artificial Sequence
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<400> 14		8
<210> 15 <211> 20 <212> DN <213> An	0	
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<400> 15		20
<210 > 16 <211 > 21 <212 > DN <213 > An	1	
	escription of Artificial Sequence: p16 reverse rimer	
<400> 16		21
<210> 17 <211> 23 <212> DN <213> An	3	
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<400> 17 acagcgto		23
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<400> 18		19

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<210> 19
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<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Control
      reverse primer
<400> 19
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                                                                   17
<210> 20
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<223> Description of Artificial Sequence: Control probe
<400> 20
cctccatggt ggtacccagc aagg
                                                                   24
<210> 21
<211> 48
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ggatccggcc accgcggccg cacgcccaat agccctgaag actattac
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<400> 22
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accessible region

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tgcata	cgtg	ggcttccaca	ggtcgtctcc	ctccggccac	tgactaact		109			
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<223>	Description of Artificial Sequence: human VEGF									
	accessible region									
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ttgggt	tttg	ccagactcca	cagtgcatac	gtgggctcca	acaggtcctc	ttccctccca	120			
gtcact							134			